

Fig.3d.
T214A,T215L,I232A & E354K mutation

Autority (Color of the Color of the Col

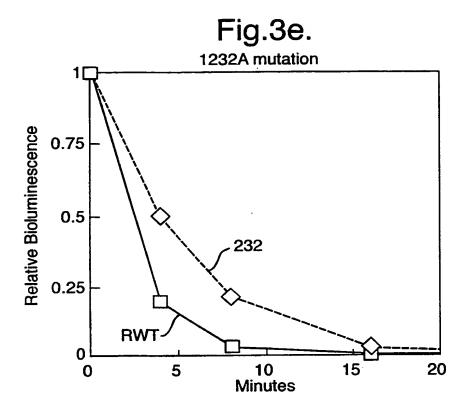


Fig.3f.

T214A & E354K mutations (44°C)

0.75

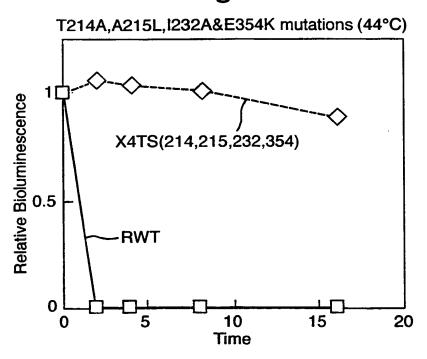
0.5

RWT

0 10

Time

Fig.3h.



## Fig.8.

CCATAAATTTACCGAATTCGTCGACTTCGATCGAGG GTGTGGAATTGTGAGCGG N-TERM.SEQ / 6651 GAGATACGCCGGGGTTCCTGG L35A-SENSE / 6652 CCAGGAACCGCGGCGTATCTC L35A-SENSE / 6653

CCCTATTTTCATTCCTGGCCAAAAGCACTG GAGTGCTTTTGGCCAGGAATGAAAATAGGG CCGCATAGAgCTCTCTGCGTCAGATTC GAATCTGACGCAGAGAGCTCTATGCGG GTTGACCGCTTGGGATCCTTAATTAAATAC F295L-SENSE/ 9048 F295L-ANTI / 9049 T214A + A215L-SENSE / 9063 T214A + A215L-ANTI / 9064 Insertion of BamHI at G339 / 9077

C-TERM.SEQ/ 6641

GTATAGATTTGAAAAAGAGCTG
CAGCTCTTTTTCAAATCTATAC
GGCTACATACTGGAGACATAGC
GCTATGTCTCCAGTATGTAGCC
GCAGTTGCGCCCGTGAACGAC
GTCGTTCACGGGCGCAACTGC
CAAATCATTCCGGGTACTGCGATTTTAAG
CAAATCATTCCGGGTACCGCGATTGTAGC
CAAATCATTCCGGGTACCGCGATTTTAG
CAAAATCGCAGTACCCGGAATGATTTG
D234G-SENSE / 792
CTTAAAAATCGCAGTACCCGGAATGATTTG
D234G-ANTI / 793

CCGCATAGAACTCTCTGCGTCAGATTC
GAATCTGACGCAGAGAGTTCTATGCGC
CTGATTACACCCAAGGGGGATG
CATCCCCCTTGGGTGTAATCAG
Cccttccgcatagannngcctgcgtcagt
CATCCCCCTTGGGTGTAATCAG
CCCTTGGGTGTAATCAG
CCCTTGGTGTAATCAG
CCCTTGGTGTAATCAG
CCCTTGGTGTAATCAG
CCCTTGGGTGTAATCAG
CCCTTGGTGTAATCAG
CCCTTGTAATCAG
CCCTTGTAATCAG
CCCTTGGTGTAATCAG
CCCTTGGTGTAATCAG
CCCTTGGTGTAATCAG
CCCTTGTAATCAG

actgacgcaggcNNNtctatgcggaaggg T214N-Anti / 82033

GCAATCAAATCGCTCCGGATACTGC 1232A-SENSE / 6911 GCAGTATCCGGAGCGATTTGATTGC 1232A-ANTI / 6912

CCAATCCATCAAGGTTTTGG H245Q-SENSE / 9128
CCAAAACCTTGATGGAATGG H245Q-ANTI / 9129